



## TRIM59 siRNA (h): sc-78105

### BACKGROUND

The tripartite motif (TRIM) family of proteins are characterized by a conserved TRIM domain that includes a coiled-coil region, a B-box type zinc finger, one RING finger and three zinc-binding domains. TRIM59 (tripartite motif-containing 59), also known as MRF1 (modulator recognition factor 1), TSBF1 or RNF104, is a 403 amino acid single-pass membrane protein that contains one RING-type zinc finger and one B box-type zinc finger. One of several members of the TRIM family, TRIM59 is thought to repress ER $\alpha$ -mediated transcriptional activation and may play a role in the regulation of ER-dependent gene expression and the pathogenesis of breast cancer. The gene encoding TRIM59 maps to human chromosome 3, which houses over 1,100 genes, including a chemo-kinase receptor (CKR) gene cluster and a variety of human cancer-related gene loci.

### REFERENCES

1. Chang, R., Xu, X. and Li, M.D. 2002. Molecular cloning, mapping and characterization of a novel mouse RING finger gene, Mrf1. *Gene* 291: 241-249.
2. Dobrzycka, K.M., Townson, S.M., Jiang, S. and Oesterreich, S. 2003. Estrogen receptor corepressors—a role in human breast cancer? *Endocr. Relat. Cancer* 10: 517-536.
3. Georgescu, S.P., Li, J.H., Lu, Q., Karas, R.H., Brown, M. and Mendelsohn, M.E. 2005. Modulator recognition factor 1, an AT-rich interaction domain family member, is a novel corepressor for estrogen receptor  $\alpha$ . *Mol. Endocrinol.* 19: 2491-2501.
4. Kominami, R. and Niwa, O. 2006. Radiation carcinogenesis in mouse thymic lymphomas. *Cancer Sci.* 97: 575-581.
5. Girault, I., Bièche, I. and Lidereau, R. 2006. Role of estrogen receptor  $\alpha$  transcriptional coregulators in tamoxifen resistance in breast cancer. *Maturitas* 54: 342-351.
6. Huang, W., Kane, J.K. and Li, M.D. 2008. Identification and characterization of a long isoform of human IFT80, IFT80-L. *Biochem. Biophys. Res. Commun.* 373: 653-658.

### CHROMOSOMAL LOCATION

Genetic locus: TRIM59 (human) mapping to 3q26.1.

### PRODUCT

TRIM59 siRNA (h) is a pool of 3 target-specific 19-25 nt siRNAs designed to knock down gene expression. Each vial contains 3 nmol of lyophilized siRNA, sufficient for a 10  $\mu$ M solution once resuspended using protocol below. Suitable for 50-100 transfections. Also see TRIM59 shRNA Plasmid (h): sc-78105-SH and TRIM59 shRNA (h) Lentiviral Particles: sc-78105-V as alternate gene silencing products.

For independent verification of TRIM59 (h) gene silencing results, we also provide the individual siRNA duplex components. Each is available as 3 nmol of lyophilized siRNA. These include: sc-78105A, sc-78105B and sc-78105C.

### RESEARCH USE

For research use only, not for use in diagnostic procedures.

### STORAGE AND RESUSPENSION

Store lyophilized siRNA duplex at -20° C with desiccant. Stable for at least one year from the date of shipment. Once resuspended, store at -20° C, avoid contact with RNases and repeated freeze thaw cycles.

Resuspend lyophilized siRNA duplex in 330  $\mu$ l of the RNase-free water provided. Resuspension of the siRNA duplex in 330  $\mu$ l of RNase-free water makes a 10  $\mu$ M solution in a 10  $\mu$ M Tris-HCl, pH 8.0, 20 mM NaCl, 1 mM EDTA buffered solution.

### APPLICATIONS

TRIM59 siRNA (h) is recommended for the inhibition of TRIM59 expression in human cells.

### SUPPORT REAGENTS

For optimal siRNA transfection efficiency, Santa Cruz Biotechnology's siRNA Transfection Reagent: sc-29528 (0.3 ml), siRNA Transfection Medium: sc-36868 (20 ml) and siRNA Dilution Buffer: sc-29527 (1.5 ml) are recommended. Control siRNAs or Fluorescein Conjugated Control siRNAs are available as 10  $\mu$ M in 66  $\mu$ l. Each contain a scrambled sequence that will not lead to the specific degradation of any known cellular mRNA. Fluorescein Conjugated Control siRNAs include: sc-36869, sc-44239, sc-44240 and sc-44241. Control siRNAs include: sc-37007, sc-44230, sc-44231, sc-44232, sc-44233, sc-44234, sc-44235, sc-44236, sc-44237 and sc-44238.

### RT-PCR REAGENTS

Semi-quantitative RT-PCR may be performed to monitor TRIM59 gene expression knockdown using RT-PCR Primer: TRIM59 (h)-PR: sc-78105-PR (20  $\mu$ l). Annealing temperature for the primers should be 55-60° C and the extension temperature should be 68-72° C.

### PROTOCOLS

See our web site at [www.scbt.com](http://www.scbt.com) for detailed protocols and support products.